T. STRZELECKA

RAW SEQUENCE LISTING

DATE: 05/31/2001

PATENT APPLICATION: US/09/675,828

TIME: 11:01:16

Input Set : C:\PAOLA\09675828.txt

Output Set: C:\CRF3\05312001\1675828.raw

SEQUENCE LISTING

```
(1) GENERAL INFORMATION:
              (i) APPLICANT: Thomas J. Cummins, Susan M. Atwood
      5
                             Lynn Bergmeyer, John B. Findlay
      6
                             John W.H. Sutherland, JoAnne H. Kerschner
      8
             (ii) TITLE OF INVENTION: DIAGNOSTIC COMPOSITIONS, ELEMENTS,
      9
                                      METHODS AND TEST KITS FOR
     10
                                      AMPLIFICATION AND DETECTION OF TWO
     11
                                      OR MORE TARGET DNA'S USING PRIMERS
W--> 12
                                      HAVING MATCHED MELTING TEMPERATURES
     14
            (iii) NUMBER OF SEQUENCES: 65
     16
             (iv) CORRESPONDENCE ADDRESS:
     17
                   (A) ADDRESSEE: Eastman Kodak Company, Patent Legal Staff
     18
                   (B) STREET: 343 State Street
     19
                   (C) CITY: Rochester
     20
                   (D) STATE: New York
                                                          ENTERED
     21
                   (E) COUNTRY: U.S.A.
     22
                   (F) ZIP: 14650 - 2201
     23
             (v) COMPUTER READABLE FORM:
     24
                   (A) MEDIUM TYPE: Diskette, 3.5inch, 1.44 MB storage (IBM)
     25
                   (B) COMPUTER: IBM PS/2
     26
                   (C) OPERATING SYSTEM: MS-DOS Version 3.3
     27
                   (D) SOFTWARE: PC-8 (Word for Windows)
     28
            (vi) CURRENT APPLICATION DATA:
C--> 29
                   (A) APPLICATION NUMBER: US/09/675,828
C--> 30
                   (B) FILING DATE: 29-Sep-2000
     31
                   (C) CLASSIFICATION:
     32
           (vii) PRIOR APPLICATION DATA:
     33
                   (A) APPLICATION NUMBER: 08/062,023
     34
                   (B) FILING DATE:
     36
          (viii) ATTORNEY/AGENT INFORMATION:
     37
                   (A) NAME: Tucker, J. Lanny
     38
                   (B) REGISTRATION NUMBER: 27,678
     39
                   (C) REFERENCE/DOCKET NUMBER: 67271A
     40
            (ix) TELECOMMUNICATION INFORMATION:
     41
                   (A) TELEPHONE: (716) 722-9332
     42
                   (B) TELEFAX: (716) 477-4646
     44
        (2) INFORMATION FOR SEQ ID NO: 1:
     45
             (i) SEQUENCE CHARACTERISTICS:
     46
                   (A) LENGTH: 28 nucleotides
     47
                   (B) TYPE: Nucleic acid
     48
                  (C) STRANDEDNESS: Single
     49
                  (D) TOPOLOGY: Linear
W--> 50
            (ii) MOLECULE TYPE: Primer for HIV-I DNA
     51
           (iii) HYPOTHETICAL: No
     52
           (iv) ANTI-SENSE: No
     53
            (vi) ORIGINAL SOURCE:
```

RAW SEQUENCE LISTING

DATE: 05/31/2001 PATENT APPLICATION: US/09/675,828 TIME: 11:01:16

Input Set : C:\PAOLA\09675828.txt

```
54
           (vii) IMMEDIATE SOURCE:
     55
             (x) PUBLICATION INFORMATION:
     56
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
     58 AGTGGGGGA CATCAAGCAG CCATGCAA 28
     62 (2) INFORMATION FOR SEQ ID NO: 2:
     63
              (i) SEQUENCE CHARACTERISTICS:
     64
                   (A) LENGTH: 28 nucleotides
     65
                   (B) TYPE: Nucleic acid
     66
                   (C) STRANDEDNESS: Single
     67
                   (D) TOPOLOGY: Linear
W--> 68
            (ii) MOLECULE TYPE: Primer for HIV-I DNA
     69
           (iii) HYPOTHETICAL: No
     70
            (iv) ANTI-SENSE: No
     71
            (vi) ORIGINAL SOURCE:
     72
           (vii) IMMEDIATE SOURCE:
     73
             (x) PUBLICATION INFORMATION:
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
     76 TTCCTGCTAT GTCACTTCCC CTTGGTTC 28
     78 (2) INFORMATION FOR SEQ ID NO: 3:
     79
             (i) SEQUENCE CHARACTERISTICS:
     80
                   (A) LENGTH: 28 nucleotides
     81
                   (B) TYPE: Nucleic acid
     82
                   (C) STRANDEDNESS: Single
     83
                   (D) TOPOLOGY: Linear
            (ii) MOLECULE TYPE: Primer for HIV-I DNA
W--> 84
     85
           (iii) HYPOTHETICAL: No
     86
            (iv) ANTI-SENSE: No
     87
            (vi) ORIGINAL SOURCE:
     88
           (vii) IMMEDIATE SOURCE:
     89
             (x) PUBLICATION INFORMATION:
     90
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
     92 TAGCACCCAC CAGGGCAAAG AGAAGAGT 28
     96 (2) INFORMATION FOR SEQ ID NO: 4:
             (i) SEQUENCE CHARACTERISTICS:
     98
                  (A) LENGTH: 28 nucleotides
     99
                   (B) TYPE: Nucleic acid
     100
                    (C) STRANDEDNESS: Single
     101
                    (D) TOPOLOGY: Linear
W--> 102
             (ii) MOLECULE TYPE: Primer for HIV-I DNA
     103
            (iii) HYPOTHETICAL: No
     104
             (iv) ANTI-SENSE: No
     105
             (vi) ORIGINAL SOURCE:
     106
            (vii) IMMEDIATE SOURCE:
     107
              (x) PUBLICATION INFORMATION:
     108
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
     110 AGATGCTGTT GCGCCTCAAT AGCCCTCA 28
     112 (2) INFORMATION FOR SEQ ID NO: 5:
     113
              (i) SEQUENCE CHARACTERISTICS:
     114
                    (A) LENGTH: 26 nucleotides
```

RAW SEQUENCE LISTING

DATE: 05/31/2001

PATENT APPLICATION: US/09/675,828

TIME: 11:01:16

Input Set : C:\PAOLA\09675828.txt

Output Set: C:\CRF3\05312001\I675828.raw

```
(B) TYPE: Nucleic acid
     115
                    (C) STRANDEDNESS: Single
     116
                    (D) TOPOLOGY: Linear
     117
             (ii) MOLECULE TYPE: Primer for HIV-I DNA
W--> 118
            (iii) HYPOTHETICAL: No
     119
             (iv) ANTI-SENSE: No
     120
             (vi) ORIGINAL SOURCE:
     121
     122
            (vii) IMMEDIATE SOURCE:
              (x) PUBLICATION INFORMATION:
     123
     124
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
     126 CTTGGTTCTC TCATCTGGCC TGGTGC 26
     130 (2) INFORMATION FOR SEQ ID NO: 6:
              (i) SEQUENCE CHARACTERISTICS:
     131
                    (A) LENGTH: 28 nucleotides
     132
                    (B) TYPE: Nucleic acid
     133
     134
                    (C) STRANDEDNESS: Single
     135
                    (D) TOPOLOGY: Linear
            (ii) MOLECULE TYPE: Probe for HIV-I DNA
W--> 136
     137
            (iii) HYPOTHETICAL: No
     138
             (iv) ANTI-SENSE: No
     139
             (vi) ORIGINAL SOURCE:
            (vii) IMMEDIATE SOURCE:
     140
              (x) PUBLICATION INFORMATION:
     141
     142
              (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
           GAGACCATCA ATGAGGAAGC TGCAGAAT
     144
     146 (2) INFORMATION FOR SEQ ID NO: 7:
              (i) SEQUENCE CHARACTERISTICS:
     147
                    (A) LENGTH: 28 nucleotides
     148
                    (B) TYPE: Nucleic acid
     149
                    (C) STRANDEDNESS: Single
     150
                    (D) TOPOLOGY: Linear
     151
             (ii) MOLECULE TYPE: Probe for HIV-I DNA
W--> 152
            (iii) HYPOTHETICAL: No
     153
     154
             (iv) ANTI-SENSE: No
     155
              (vi) ORIGINAL SOURCE:
             (vii) IMMEDIATE SOURCE:
     156
              (x) PUBLICATION INFORMATION:
     157
              (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
     158
     160 GTGCAGCAGC AGAACAATTT GCTGAGGG 28
     164 (2) INFORMATION FOR SEQ ID NO: 8:
     165
              (i) SEQUENCE CHARACTERISTICS:
                    (A) LENGTH: 30 nucleotidses
     166
     167
                    (B) TYPE: Nucleic acid
                    (C) STRANDEDNESS: Single
     168
                    (D) TOPOLOGY: Linear
     169
W--> 170
             (ii) MOLECULE TYPE: Nonsense probe
```

(iii) HYPOTHETICAL: No

(vi) ORIGINAL SOURCE:

(iv) ANTI-SENSE: No

171

172

173

RAW SEQUENCE LISTING DATE: 05/31/2001 PATENT APPLICATION: US/09/675,828 TIME: 11:01:16

Input Set : C:\PAOLA\09675828.txt

```
(vii) IMMEDIATE SOURCE:
     174
              (x) PUBLICATION INFORMATION:
     175
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
     176
     178 GGTGTCACCC CCAGAGTCCC CTGTACCCGC
     180 (2) INFORMATION FOR SEQ ID NO: 9:
              (i) SEQUENCE CHARACTERISTICS:
     181
                   (A) LENGTH: 41 nucleotides
     182
     183
                   (B) TYPE: Nucleic acid
                   (C) STRANDEDNESS: Single
     184
     185
                   (D) TOPOLOGY: Linear
             (ii) MOLECULE TYPE: Oligonucleotide from HIV-I DNA
W--> 186
     187
            (iii) HYPOTHETICAL: No
     188
             (iv) ANTI-SENSE: No
     189
             (vi) ORIGINAL SOURCE:
     190
            (vii) IMMEDIATE SOURCE:
     191
              (x) PUBLICATION INFORMATION:
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
     192
     194 ATCCTGGGAT TAAATAAAAT AGTAAGAATG TATAGCCCTA C
     197 (2) INFORMATION FOR SEQ ID NO: 10:
              (i) SEQUENCE CHARACTERISTICS:
                    (A) LENGTH: 25 nucleotides
     199
     200
                    (B) TYPE: Nucleic acid
                    (C) STRANDEDNESS: Single
     201
     202
                    (D) TOPOLOGY: Linear
             (ii) MOLECULE TYPE: Primer for hCMV DNA
W--> 203
            (iii) HYPOTHETICAL: No
     204
             (iv) ANTI-SENSE: No
             (vi) ORIGINAL SOURCE:
     206
     207
            (vii) IMMEDIATE SOURCE:
              (x) PUBLICATION INFORMATION:
     208
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
     209
     211 TGCACTGCCA GGTGCTTCGG CTCAT
     213 (2) INFORMATION FOR SEQ ID NO: 11:
              (i) SEQUENCE CHARACTERISTICS:
     214
     215
                    (A) LENGTH: 25 nucleotides
     216
                   (B) TYPE: Nucleic acid
     217
                    (C) STRANDEDNESS: Single
     218
                    (D) TOPOLOGY: Linear
W--> 219
             (ii) MOLECULE TYPE: Primer for hCMV DNA
     220
           (iii) HYPOTHETICAL: No
     221
             (iv) ANTI-SENSE: No
     222
             (vi) ORIGINAL SOURCE:
     223
            (vii) IMMEDIATE SOURCE:
              (x) PUBLICATION INFORMATION:
     224
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
     225
     227 CACCACGCAG CGGCCCTTGA TGTTT
     231 (2) INFORMATION FOR SEQ ID NO: 12:
     232
              (i) SEQUENCE CHARACTERISTICS:
     233
                    (A) LENGTH: 30 nucleotides
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/675,828

DATE: 05/31/2001 TIME: 11:01:16

Input Set : C:\PAOLA\09675828.txt

```
234
                   (B) TYPE: Nucleic acid
     235
                   (C) STRANDEDNESS: Single
                   (D) TOPOLOGY: Linear
     236
W--> 237
             (ii) MOLECULE TYPE: Probe for hCMV DNA
     238
            (iii) HYPOTHETICAL: No
     239
             (iv) ANTI-SENSE: No
     240
             (vi) ORIGINAL SOURCE:
     241
            (vii) IMMEDIATE SOURCE:
     242
              (x) PUBLICATION INFORMATION:
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
     243
     245 GAACCGAGGG CCGGCTCACC TCTATGTTGG 30
     247 (2) INFORMATION FOR SEQ ID NO: 13:
             (i) SEQUENCE CHARACTERISTICS:
     248
     249
                   (A) LENGTH: 30 nucleotides
     250
                   (B) TYPE: Nucleic acid
     251
                   (C) STRANDEDNESS: Single
     252
                   (D) TOPOLOGY: Linear
W--> 253
            (ii) MOLECULE TYPE: Primer for HIV-I DNA
     254
            (iii) HYPOTHETICAL: No
            (iv) ANTI-SENSE: No
     255
     256
             (vi) ORIGINAL SOURCE:
     257
            (vii) IMMEDIATE SOURCE:
     258
              (x) PUBLICATION INFORMATION:
     259
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
     261 CCTGCTATGT CACTTCCCCT TGGTTCTCTC 30
     265 (2) INFORMATION FOR SEQ ID NO: 14:
     266
              (i) SEQUENCE CHARACTERISTICS:
     267
                   (A) LENGTH: 27 nucleotides
     268
                   (B) TYPE: Nucleic acid
     269.
                   (C) STRANDEDNESS: Single
     270
                   (D) TOPOLOGY: Linear
W--> 271
             (ii) MOLECULE TYPE: Primer for HIV-II DNA
     272
            (iii) HYPOTHETICAL: No
     273
             (iv) ANTI-SENSE: No
     274
             (vi) ORIGINAL SOURCE:
     275
            (vii) IMMEDIATE SOURCE:
     276
              (x) PUBLICATION INFORMATION:
     277
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
     279 AAGTAGACCA ACAGCACCAC CTAGCGG
     281 (2) INFORMATION FOR SEO ID NO: 15:
              (i) SEQUENCE CHARACTERISTICS:
     282
     283
                   (A) LENGTH: 29 nucleotides
     284
                   (B) TYPE: Nucleic acid
     285
                   (C) STRANDEDNESS: Single
     286
                   (D) TOPOLOGY: Linear
W--> 287
             (ii) MOLECULE TYPE: Primer for HIV-II DNA
     288
            (iii) HYPOTHETICAL: No
     289
            (iv) ANTI-SENSE: No
     290
             (vi) ORIGINAL SOURCE:
```

VERIFICATION SUMMARY

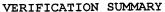
DATE: 05/31/2001 PATENT APPLICATION: US/09/675,828 TIME: 11:01:17

Input Set : C:\PAOLA\09675828.txt

Output Set: C:\CRF3\05312001\I675828.raw

L:29 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:] L:30 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:] L:12 M:259 W: Allowed number of lines exceeded, (ii) TITLE OF INVENTION: L:56 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=1 L:50 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=1 L:74 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=2 L:68 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=2 L:90 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=3 L:84 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=3 L:108 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=4 L:102 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=4 L:124 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=5 L:118 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=5 L:142 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=6 L:136 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=6 L:158 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=7 L:152 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=7 L:176 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=8 L:170 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=8 L:192 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=9
L:186 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=9 L:209 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=10 L:203 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=10 L:225 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=11 L:219 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=11 L:243 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=12 L:237 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=12 L:259 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=13 L:253 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=13 L:277 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=14 L:271 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=14 L:293 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=15 L:287 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=15 L:311 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=16 L:305 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=16 L:327 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=17 L:321 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=17 L:345 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=18 L:339 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=18 L:361 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=19 L:355 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=19 L:379 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=20 L:373 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=20 L:395 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=21 L:389 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=21 L:413 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=22 L:407 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=22 L:429 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=23

DATE: 05/31/2001



PATENT APPLICATION: US/09/675,828 TIME: 11:01:17

Input Set : C:\PAOLA\09675828.txt

```
L:423 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=23
L:447 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=24
L:441 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=24
L:463 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=25
L:457 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=25
L:481 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=26
L:475 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=26
L:497 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=27 L:491 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=27
L:515 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=28
L:509 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=28
L:531 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=29
L:525 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=29
L:549 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=30
L:543 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=30
L:565 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=31
L:559 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=31
L:583 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=32
L:577 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=32
L:599 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=33
L:593 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=33
L:617 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=34
L:611 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=34 L:633 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=35
L:627 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=35
L:651 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=36
L:645 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=36
L:667 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=37
L:661 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=37
L:685 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=38
L:679 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=38 L:701 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=39
L:695 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=39
L:720 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=40
L:713 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=40
L:737 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=41
L:730 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=41
L:756 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=42
L:749 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=42
L:773 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=43
L:766 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=43
L:792 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=44
L:785 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=44
L:809 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=45
L:802 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=45
L:827 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=46
L:821 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=46 L:843 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=47
L:837 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=47
```



DATE: 05/31/2001 PATENT APPLICATION: US/09/675,828 TIME: 11:01:17

Input Set : C:\PAOLA\09675828.txt

Output Set: C:\CRF3\05312001\I675828.raw

L:861 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=48 L:855 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=48 L:877 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=49 L:871 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=49 L:895 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=50 L:889 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=50 L:911 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=51 L:905 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=51 L:929 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=52 L:923 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=52 L:945 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=53 L:939 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=53 L:964 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=54 L:958 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=54 L:980 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=55 L:974 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=55 L:998 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=56 L:992 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=56 L:1015 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=57 L:1008 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=57 L:1034 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=58 L:1027 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=58 L:1051 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=59 L:1044 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=59 L:1070 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=60 L:1063 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=60 L:1087 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=61 L:1080 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=61 L:1105 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=62 L:1099 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=62 L:1121 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=63 L:1115 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=63 L:1139 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=64 L:1133 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=64 L:1155 M:246 W: Invalid value of Alpha Sequence Header Field, [FEATURE:], SeqNo=65 L:1149 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=65